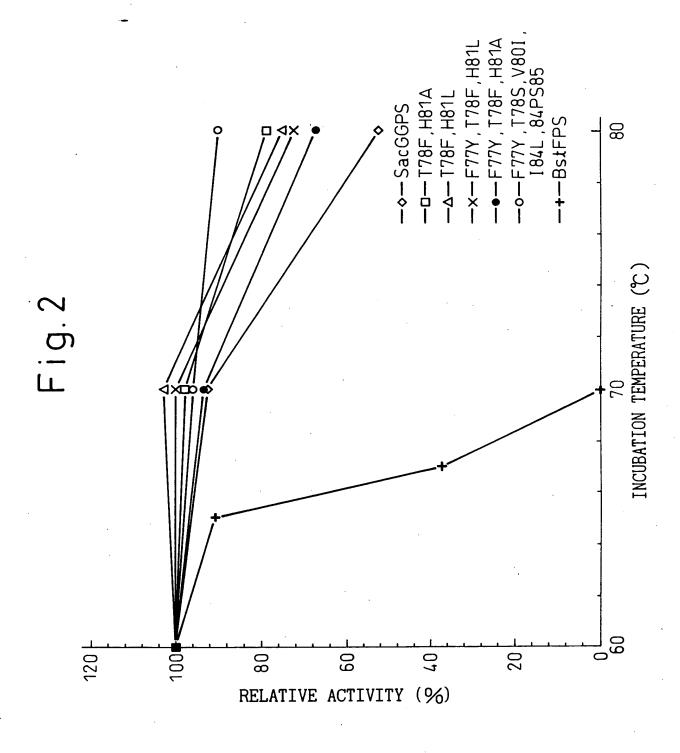
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REGION III 238 GQVVD 230	238 ··· A· 211 ·· Y··	8	313 ··AWE	Σ·· 66	9		IADK	· · · · · · · · · · · · · · · · · · ·	\	EEI.	REG.	·HAR	ANER		<u> </u>	TEG.
EMIHTMSLIHDDLPCMDNDDLRRG	A D . P V	·FT·V···I··Q·NI··· CA··V·····F·DAEI···	A	FL VA .QA	A ·· LV V ·· VE· SV ····		IL DVTKSSK ELGKTAGKDLI	. Λ	$\overline{}$	·TE·KK··YDGGAE·GMMEMAE	· · GLTADEKE · · · · PVFS · IR	TPQ	_AMPQ.		L.GLFGD.NV A.A.DG.F	AN · G
REGION I 147 EMIHT 141	149 ······ 119 ··V·AA	74 ·VL··F 95 ·L··CA	223 ·LM·CA 86 ·LT··A	36.		REGION V	GLLFQVVDDI	W	•	• M • Y • • • •	·IA··I··	• E A • • • A • •	·SA··IA··	.QALL	·· AY · LR ··	·· VI ·· I ··
REGION I GGKRVR	 PAP	L. AI.		·-i	·DI·	REGION IV	σ	∞	δ	F 256	_	\sim	\mathbf{C}	\sim	4	. 272
REC 116 G 110 •	118 88	43 64	192 · 54 ·	104	197		265 K	255	263	230	185	203	331	199	226	286
ATGERPYRS LA15778.p	CAGERDIS. ATGGPSRP.	GGPS.pep SPCRT.pep	RCPHSYNG. EHCRTS.pe	MXCRTNODA	NCAL3.pep		ATGERPYRS	LA15778.p	CAGERDIS.	ATGGPSRP.	GGPS.pep	SPCRT.pep	RCPHSYNG.	EHCRTS.pe	MXCRTNODA	NCAL3.pep

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Fig. 3

--- s.f.

___ GOH

- FOH

----GGOH

----GFOH

SacGGPS T78F, H81A

778F, H81L

F77Y, T78F, H81L F77Y, T78F, H81A ori.